ABSTRACT OF THE DISCLOSURE

The method of differentiating beer yeast of the invention is a method which comprises a first step of synthesizing a primer capable of amplifying the linker portion between a base sequence (A) and a base sequence (B) in a novel gene (C) which has the base sequence (B) comprising a portion of yeast chromosome IX linked downstream from the base sequence (A) comprising a portion of the N-terminal end of yeast gene Lg-FLO1, and which includes the base sequences listed as SEQ. ID. Nos. 1-6 of the Sequence Listing; a second step of carrying out a PCR (Polymerase Chain Reaction) using the primer synthesized in the first step and DNA separated from a yeast specimen; and a third step of differentiating whether the yeast is bottom-fermenting yeast or wild yeast, based on the PCR amplification product obtained from the second step.